

Fig. 1

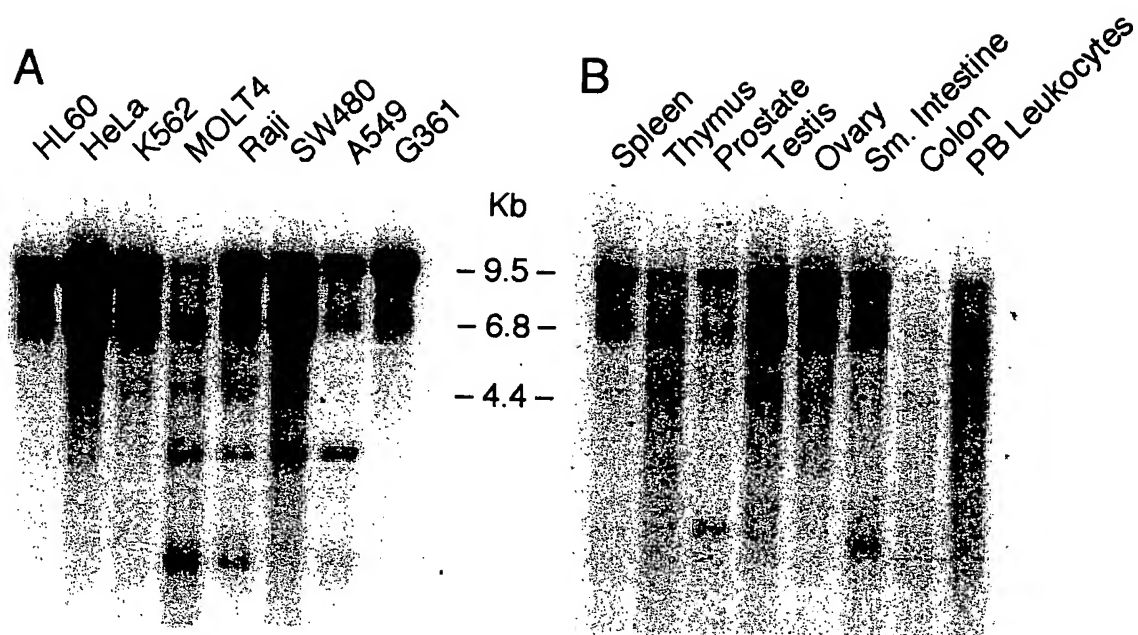


Fig. 2

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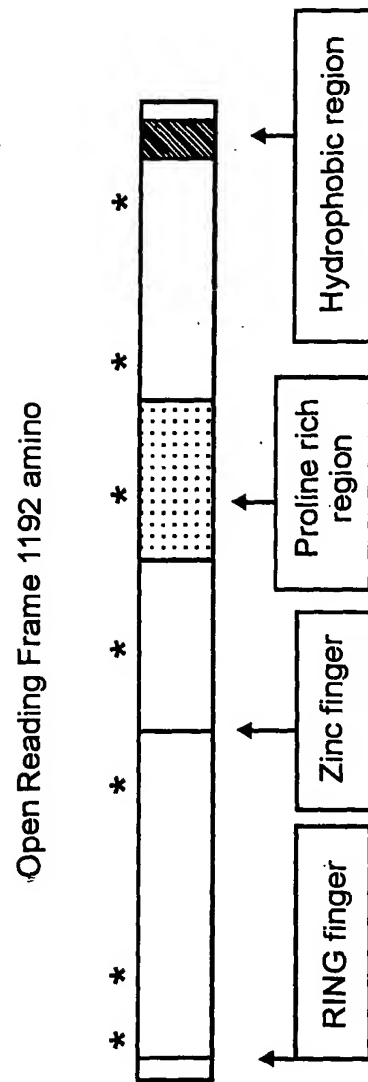


Fig. 3

A. C3HC3D Ring finger homologies

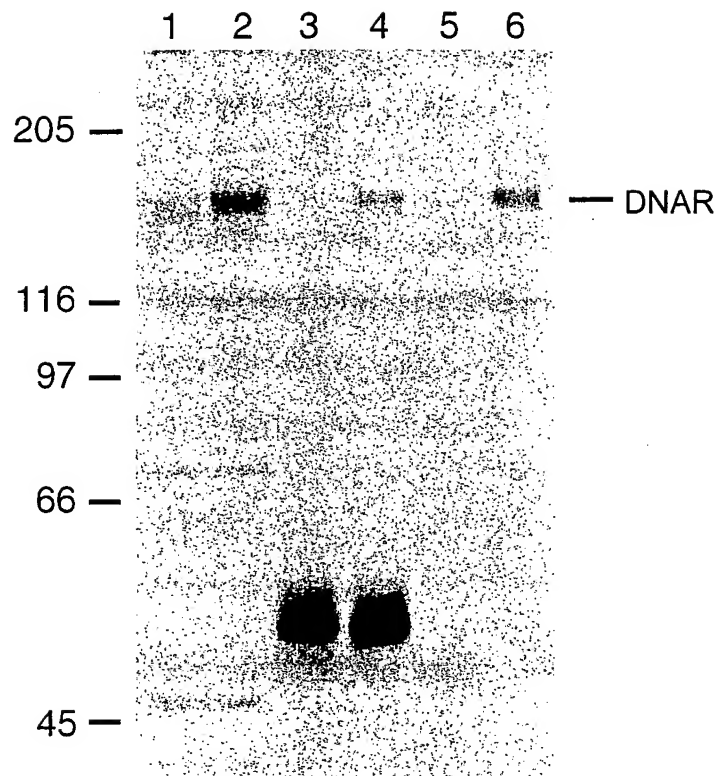
MNAB	TEFLSCPIC---	YNEFDENVHKPISLGC	SHTVCKTCLNKLHRKA-----	CPFDQTA	58			
H.sapiens ARD1	VKVL	ECGVCEDVFSL--	QGDKVPRLLLCGHTV	CHDCLTRLPLHGR--	AIRCPFDRQVTD	80		
H.sapiens CART1	KRRL	CPLCGKPMREP	VQVST-----	CGHRFCDTCLQEF	LSEG---VFKCPEDQLPLD	62		
H.sapiens SBBIO3	DEDL	ICPICSGVL	EEPVAAPH-----	CEHAFCNACITQWFS	QQ---Q--TCPVDRSVVT	61		
C.elegans cDNA EST 3879246	--	YSECLVC--	YQKFDENTRIPR	VMDCGHTLCDFCINQIV	KMAGCYSATCPFDRVRI-	182		
C.elegans 25.8KD protein	-	EWRS	CFICTMEYSR	TDKNLH-P	IIILNCGHNLCRSCINKL---	TGNGIVKCPFDR---	199	
C.elegans cDNA EST 3878739	QEV	LCCSICNRRH	FN-----	TFLPVSL	ICGHVICRKCAEK	PENQTK-----	PCPHDDWKT	61
C3HC3D motif	----	C--C-----	-----	C-H--C-----	-----	C--D-----		

B. C3H Zinc finger homologies

MNAB	ETPQPQPN	SKYKTS	MCRDLR	QQG	QCP	RGTNCT	FAHSQEE	ELEKYRLRNK	448	
C. elegans	PIE-1	-----	HTEYK	TRL	CDA	FRREGY	CPYNDNCT	YAHGQDEL	VPRRRQE	136
D.melanogaster	DTIS 11	---	QPMNTS	RYKTEL	CRP	FE	EAGECKY	GEKCCQ	FAHGSHEL	166
H.sapiens	TIS11B	-----	SSRYK	TEL	CRP	FE	ENGACKY	GDKCCQ	FAHGIHEL	181
S.cerevisiae	CTH1	QLPQLVN	KTL	YKTEL	EL	CES	FTIKGY	CKYGNK	CQFAHGLNELK	235
C3H motif	-----	YKTEL	C-----	C-----	C-----	C-----	C-----	C-----	C-----	

Fig. 4

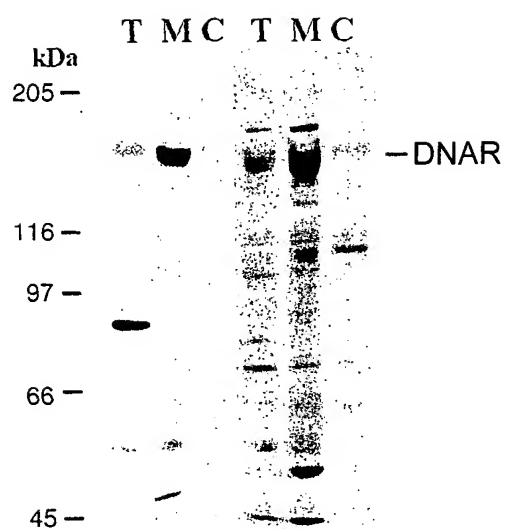
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Fig. 5

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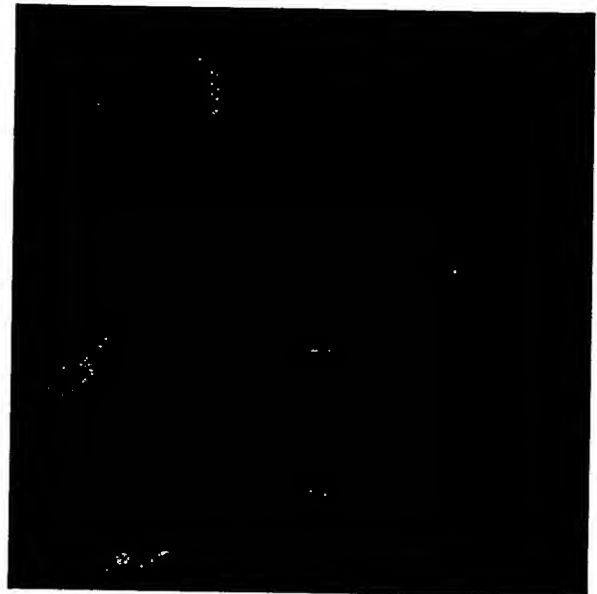
Fig. 6

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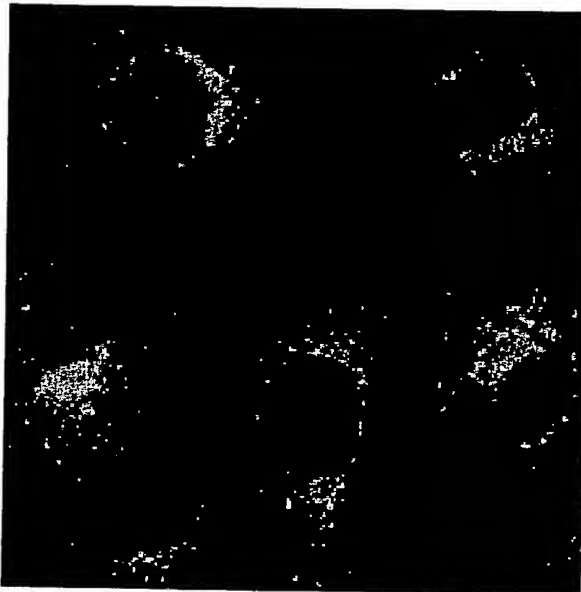
A



B



C



D



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Fig. 7

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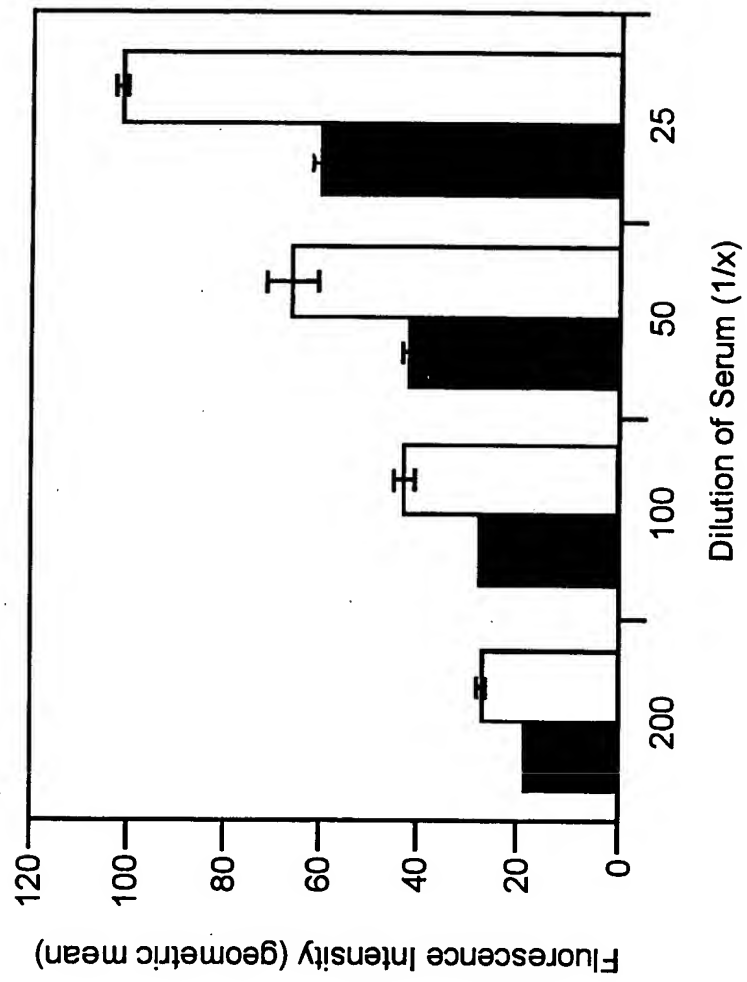


Fig. 8A

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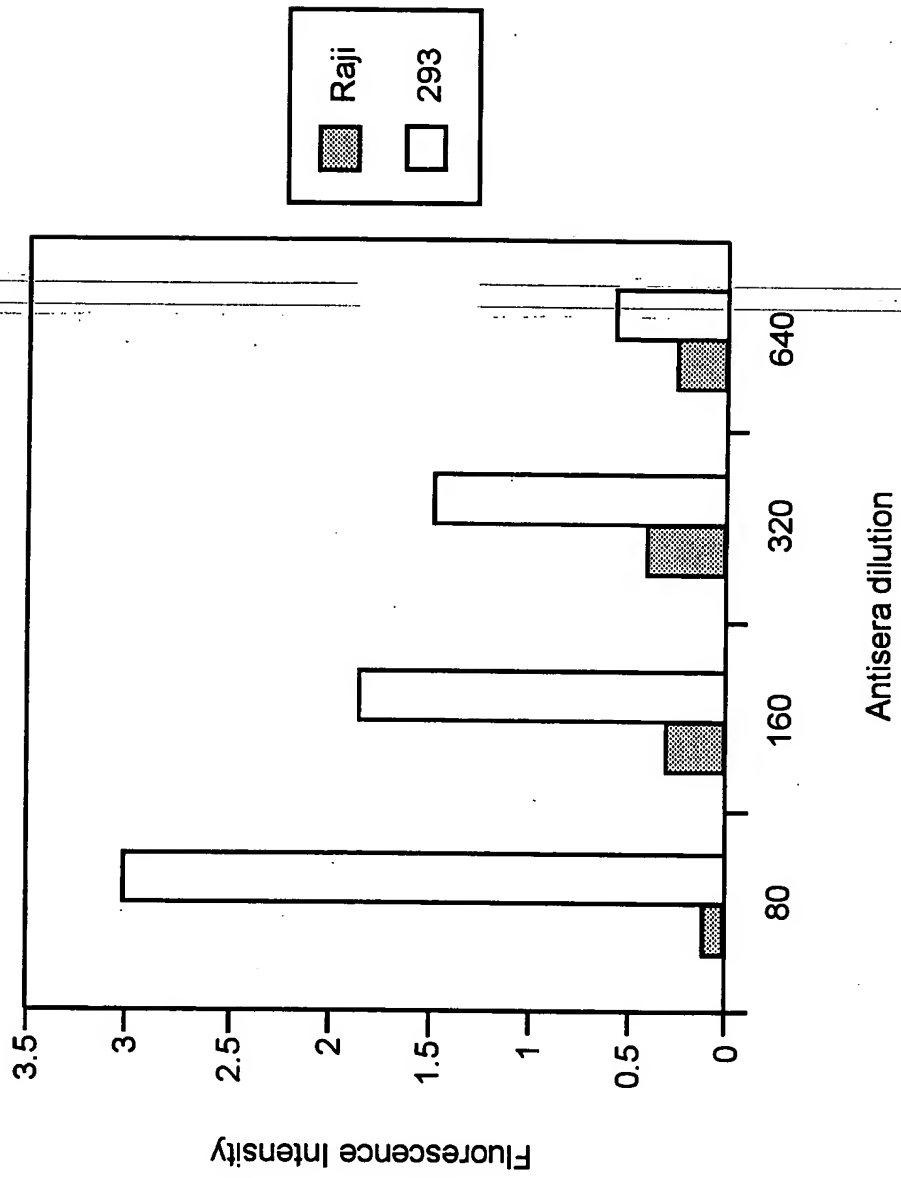
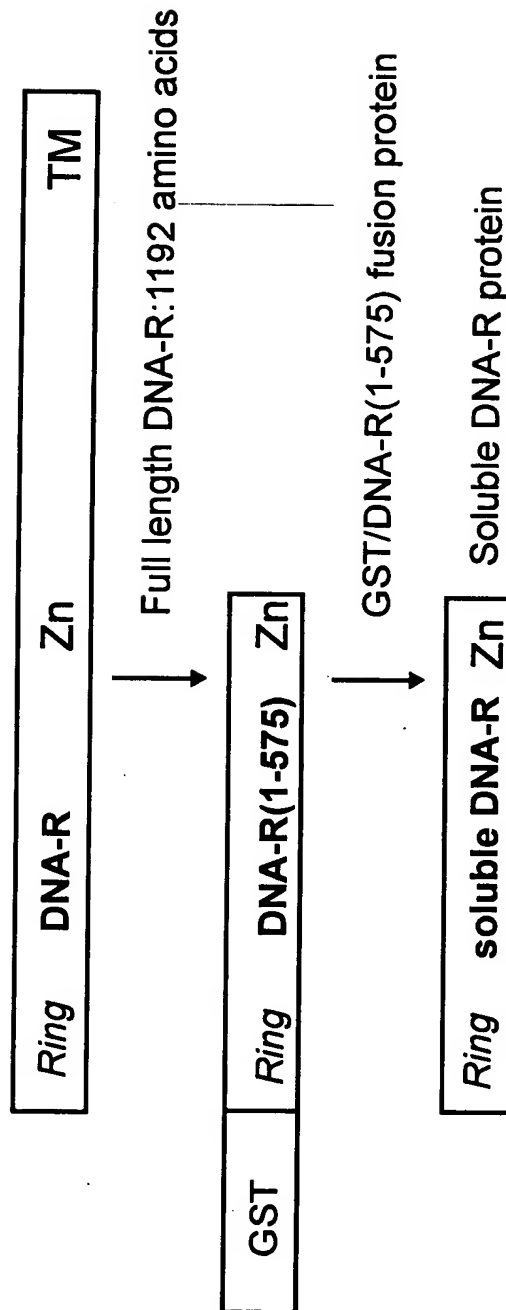
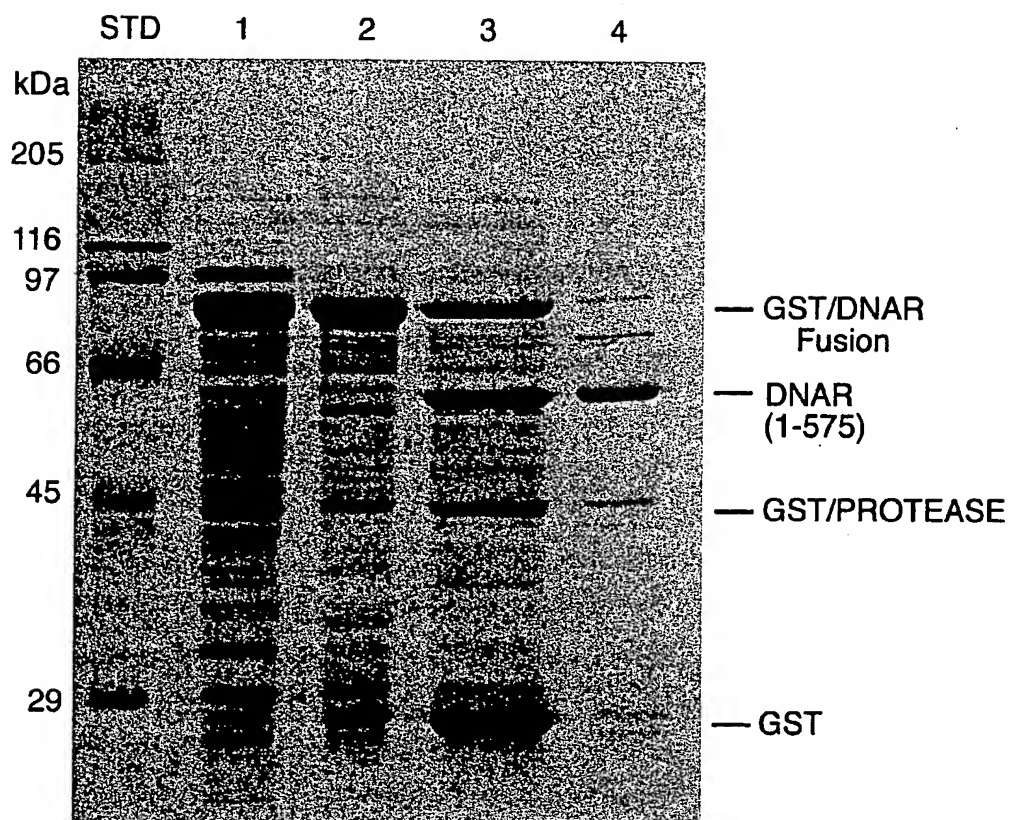


Fig. 8B

*Fig. 9A*

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Fig. 9B

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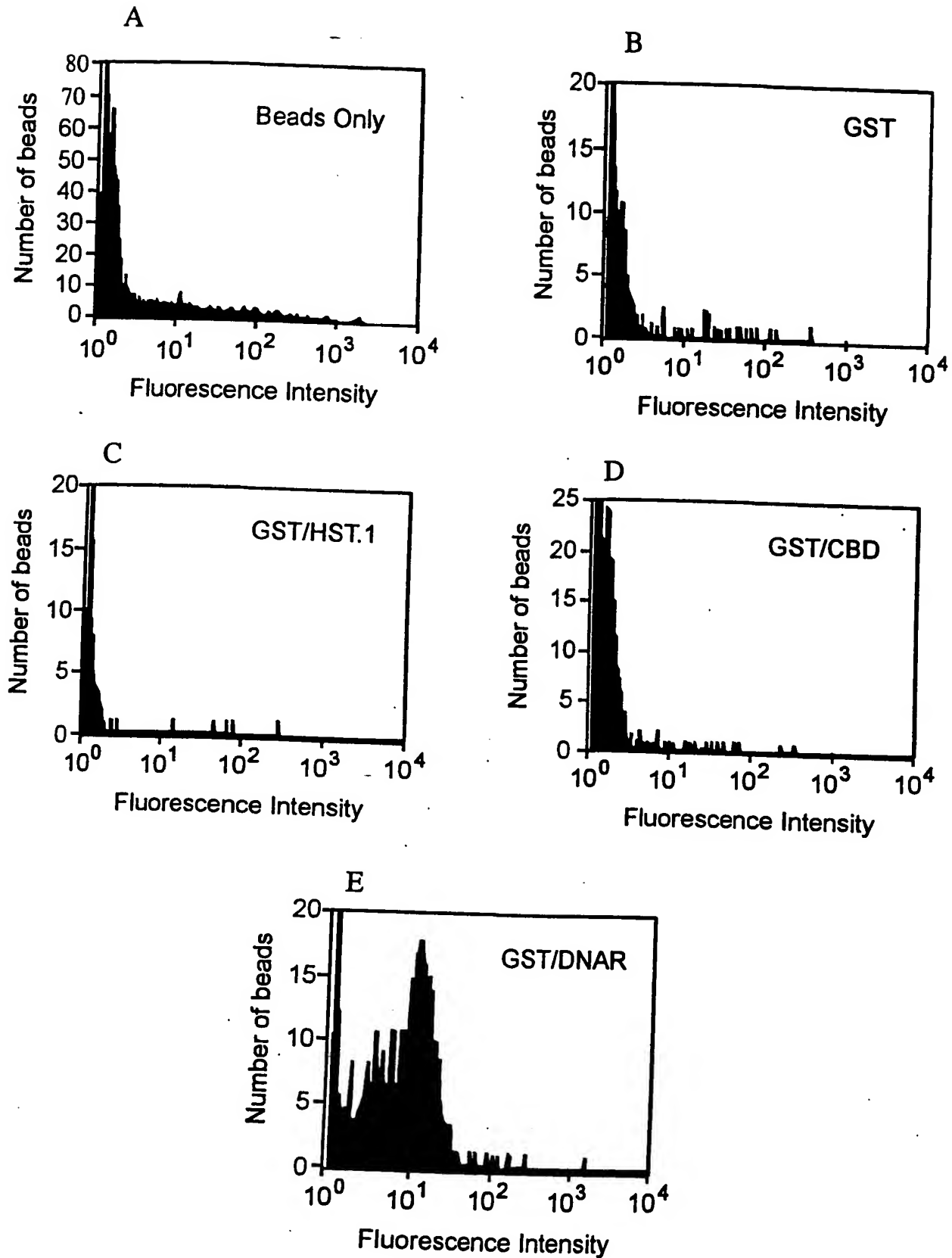


Fig. 10

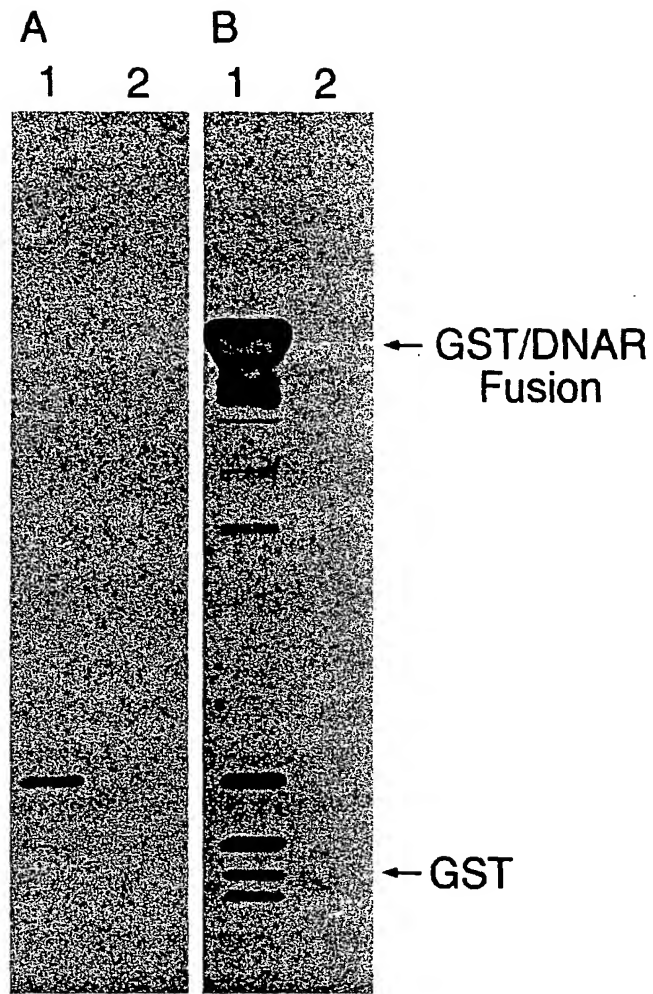


Fig. 11

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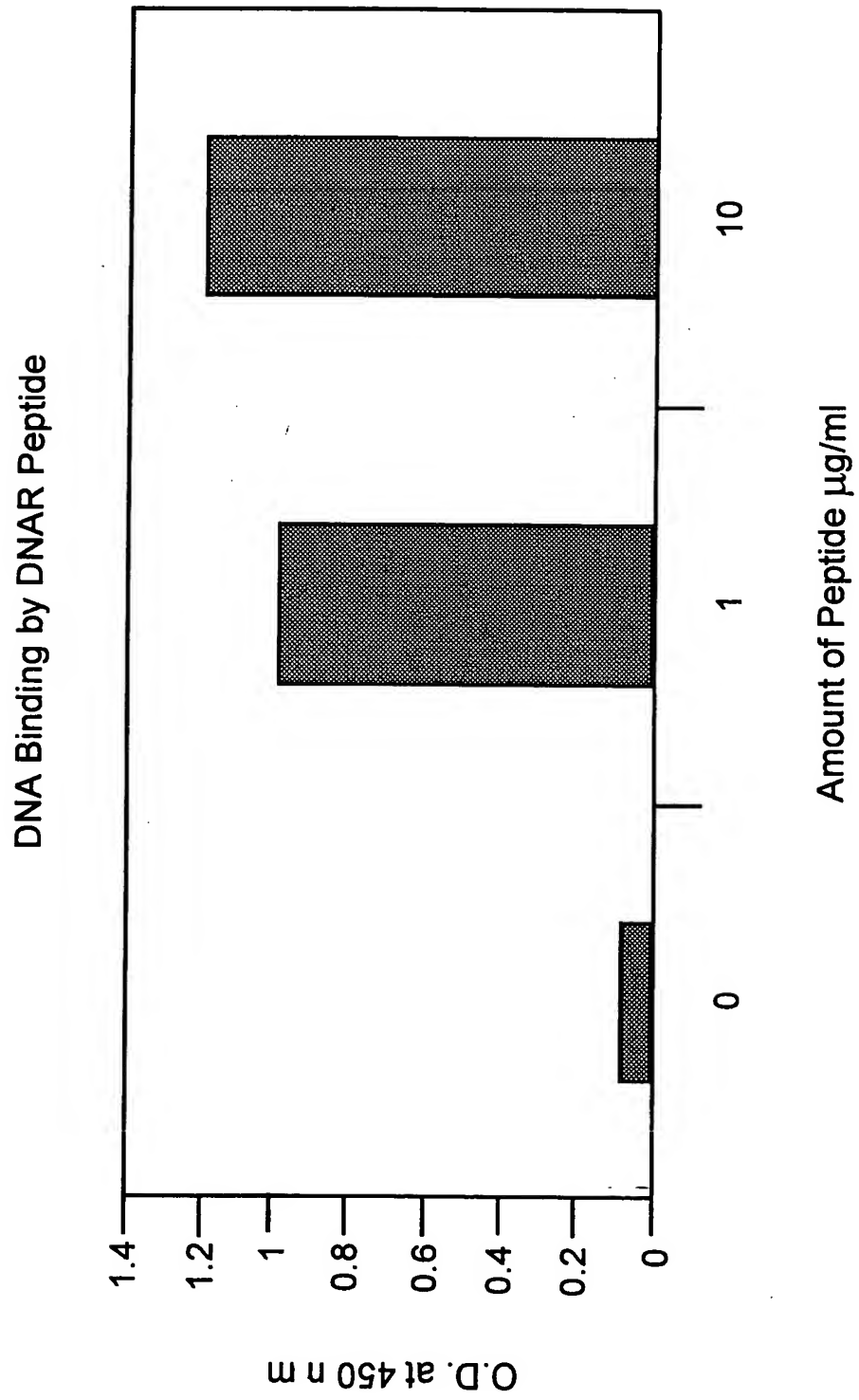
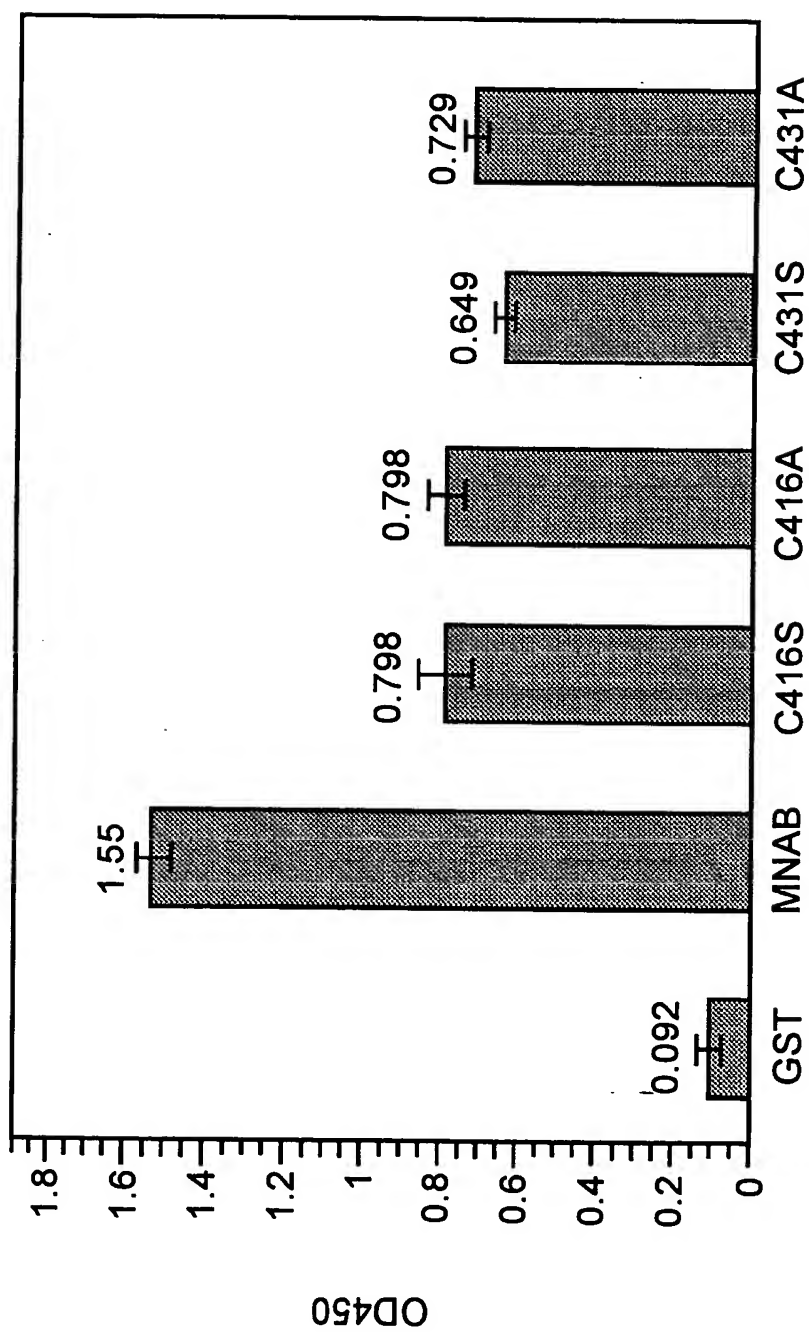
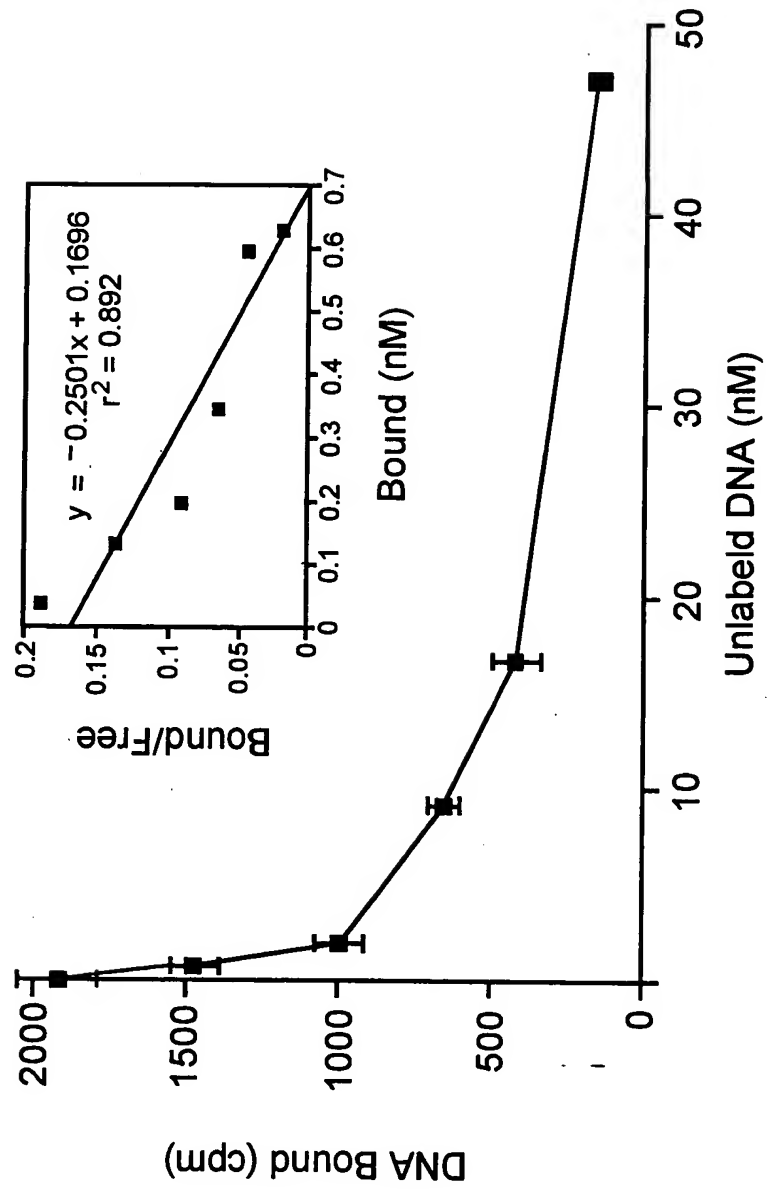
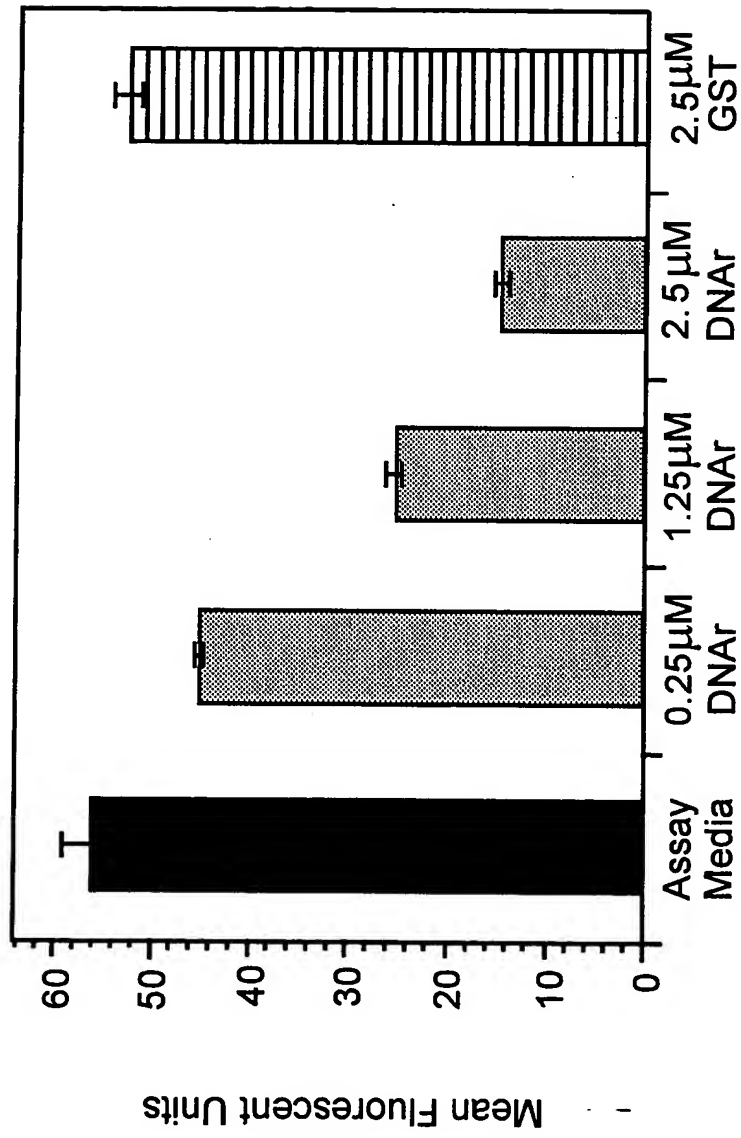


Fig. 12

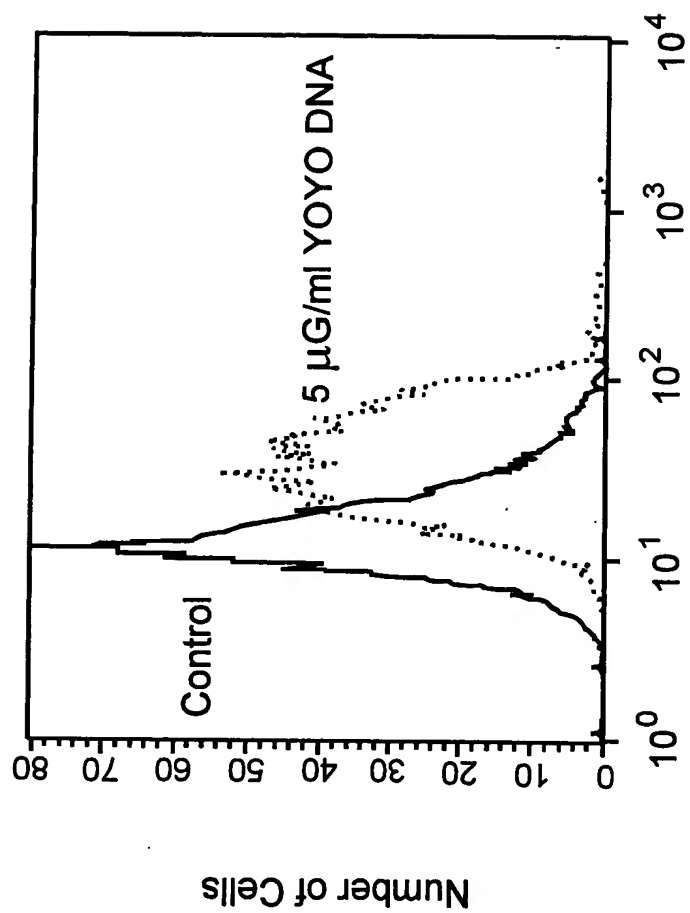
*Fig. 13*

*Fig. 14*



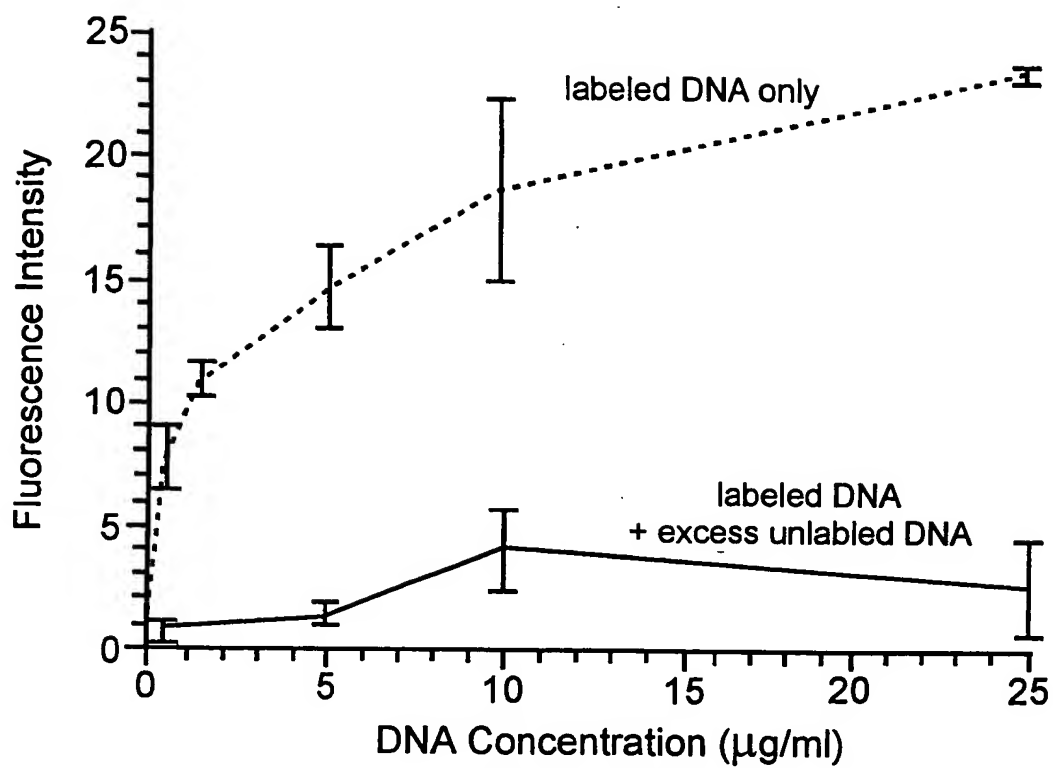
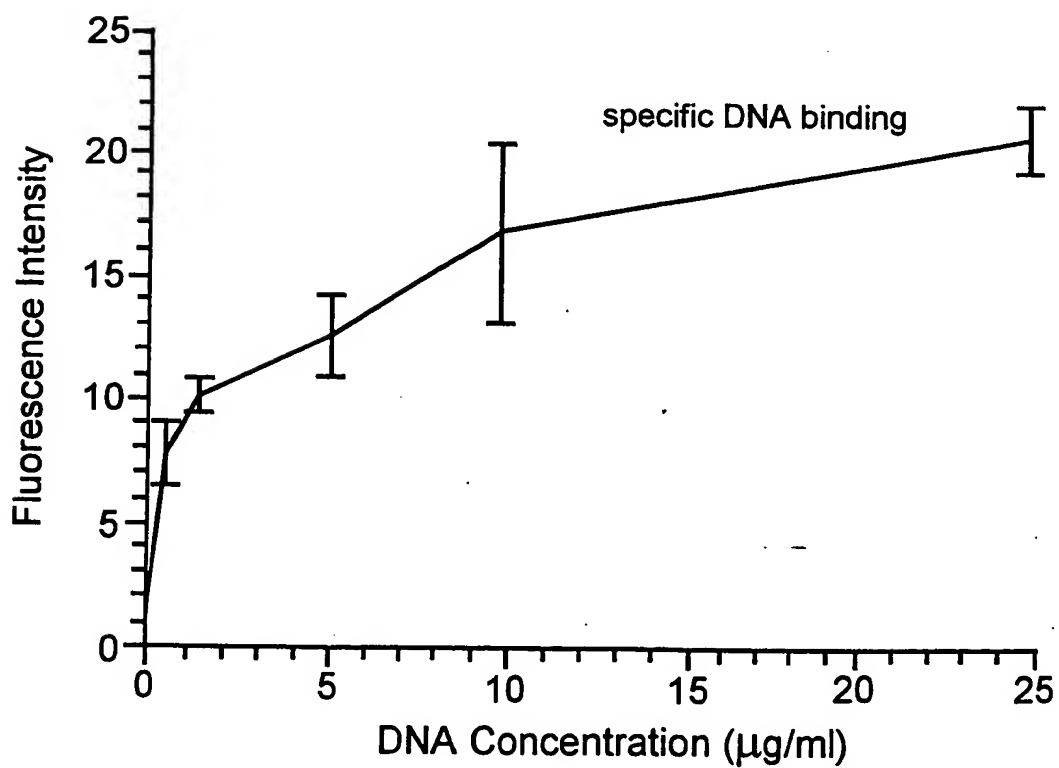
Concentration of Competitor Protein

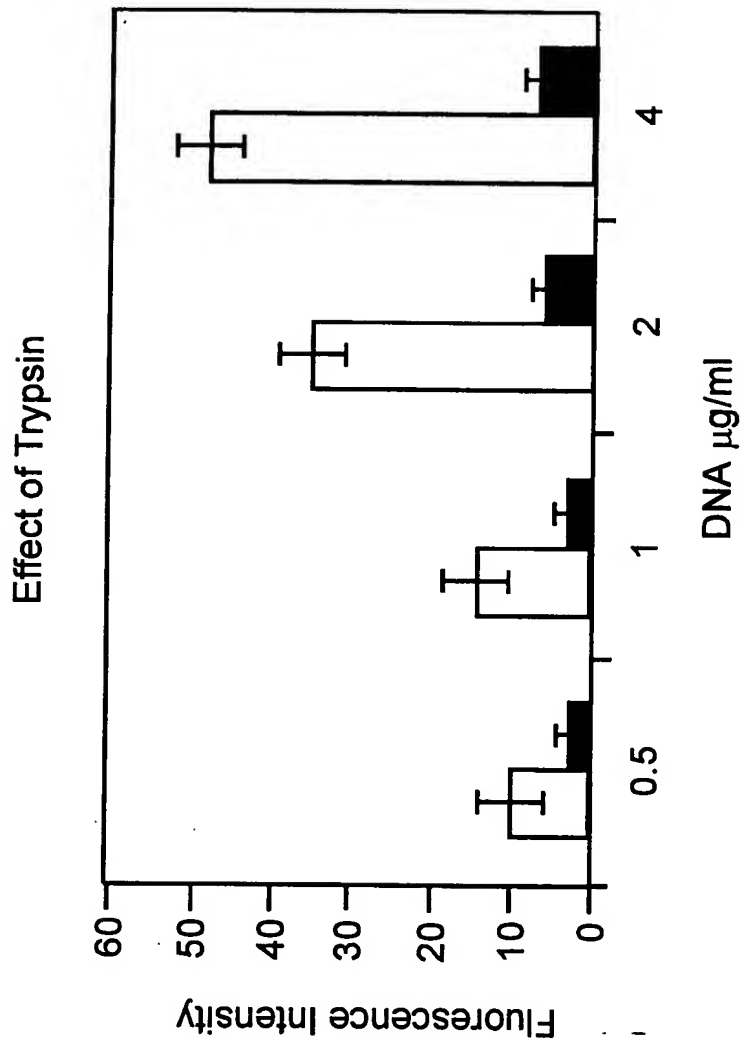
Fig. 15

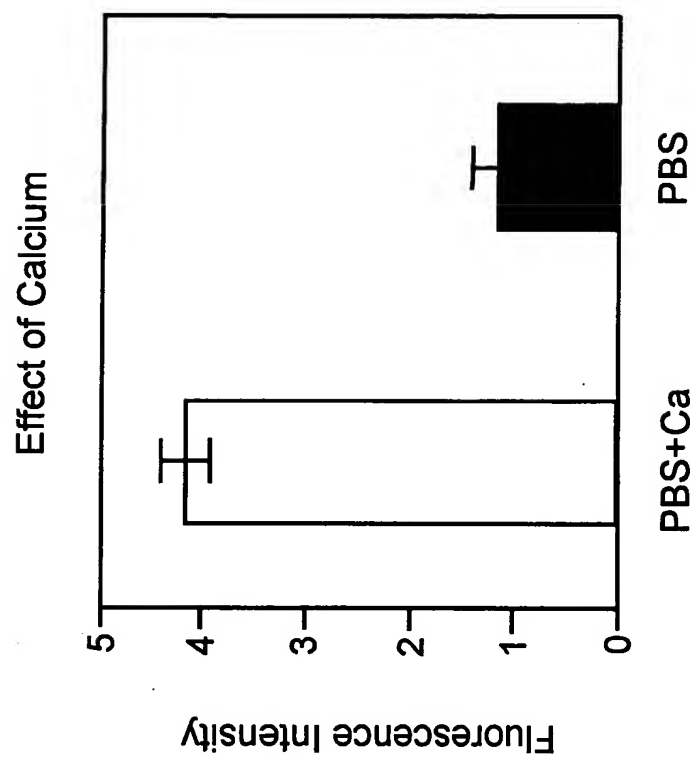


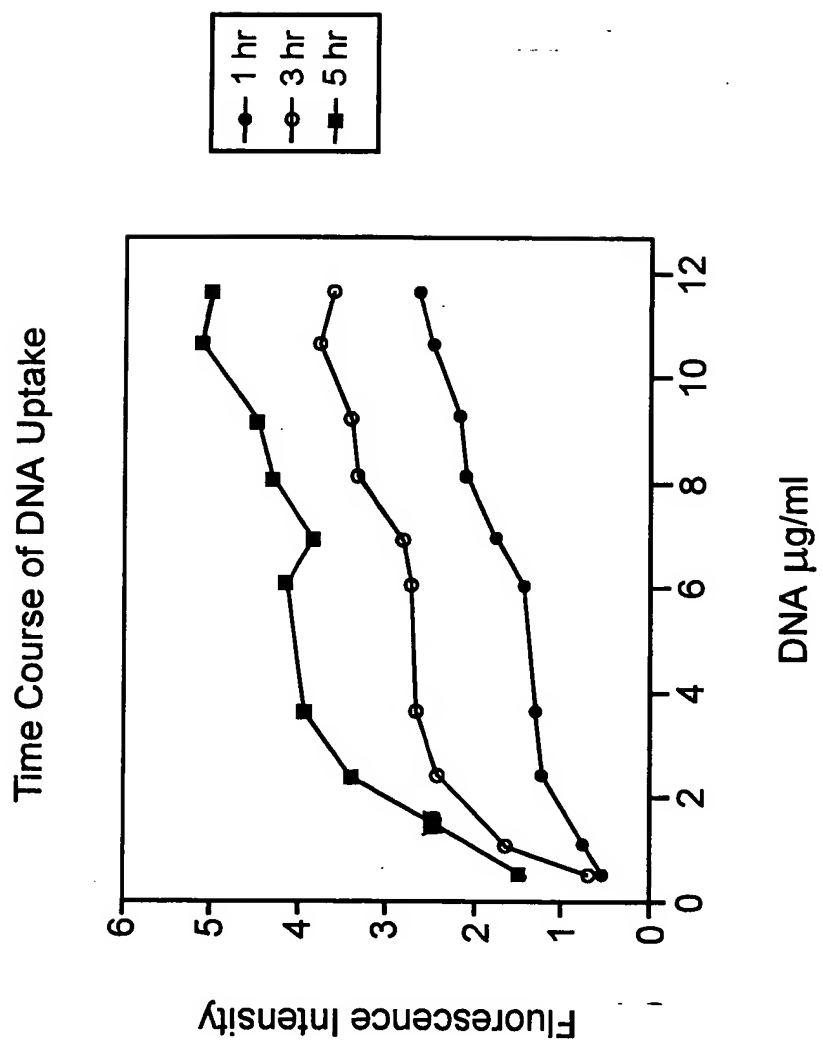
Fluorescence Intensity

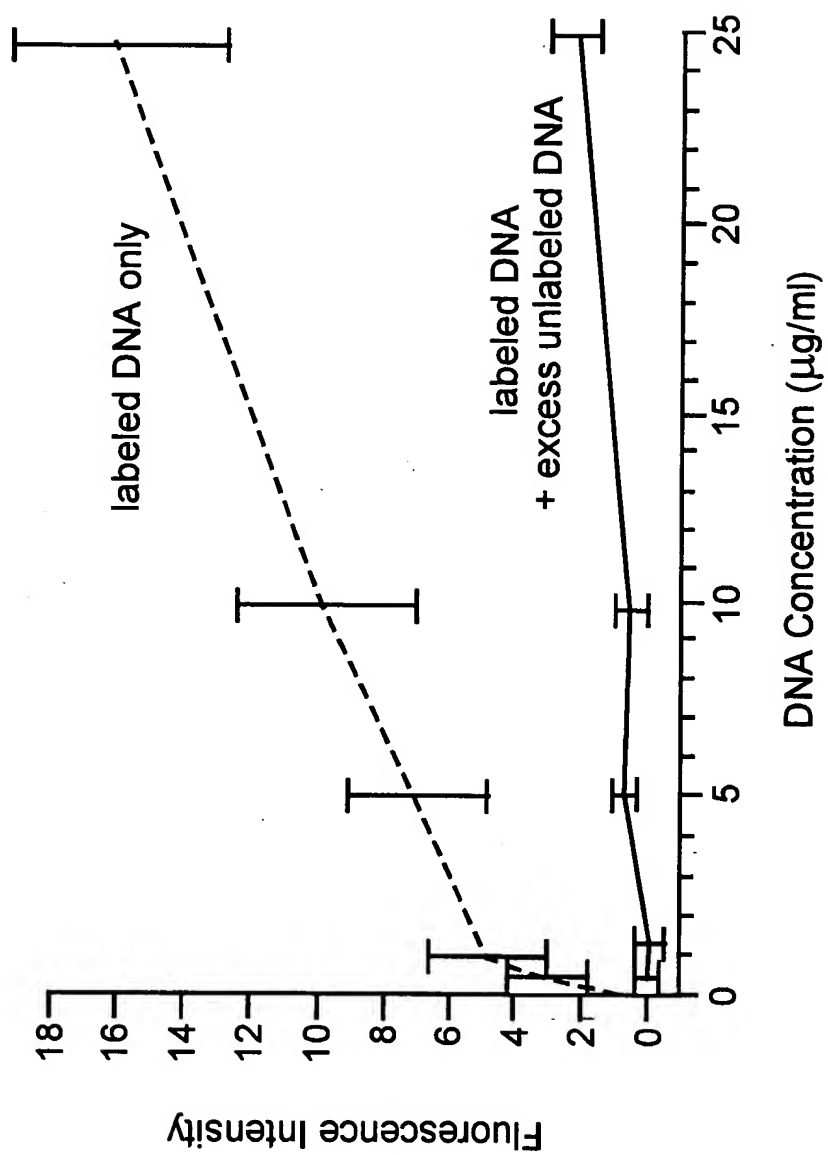
Fig. 16

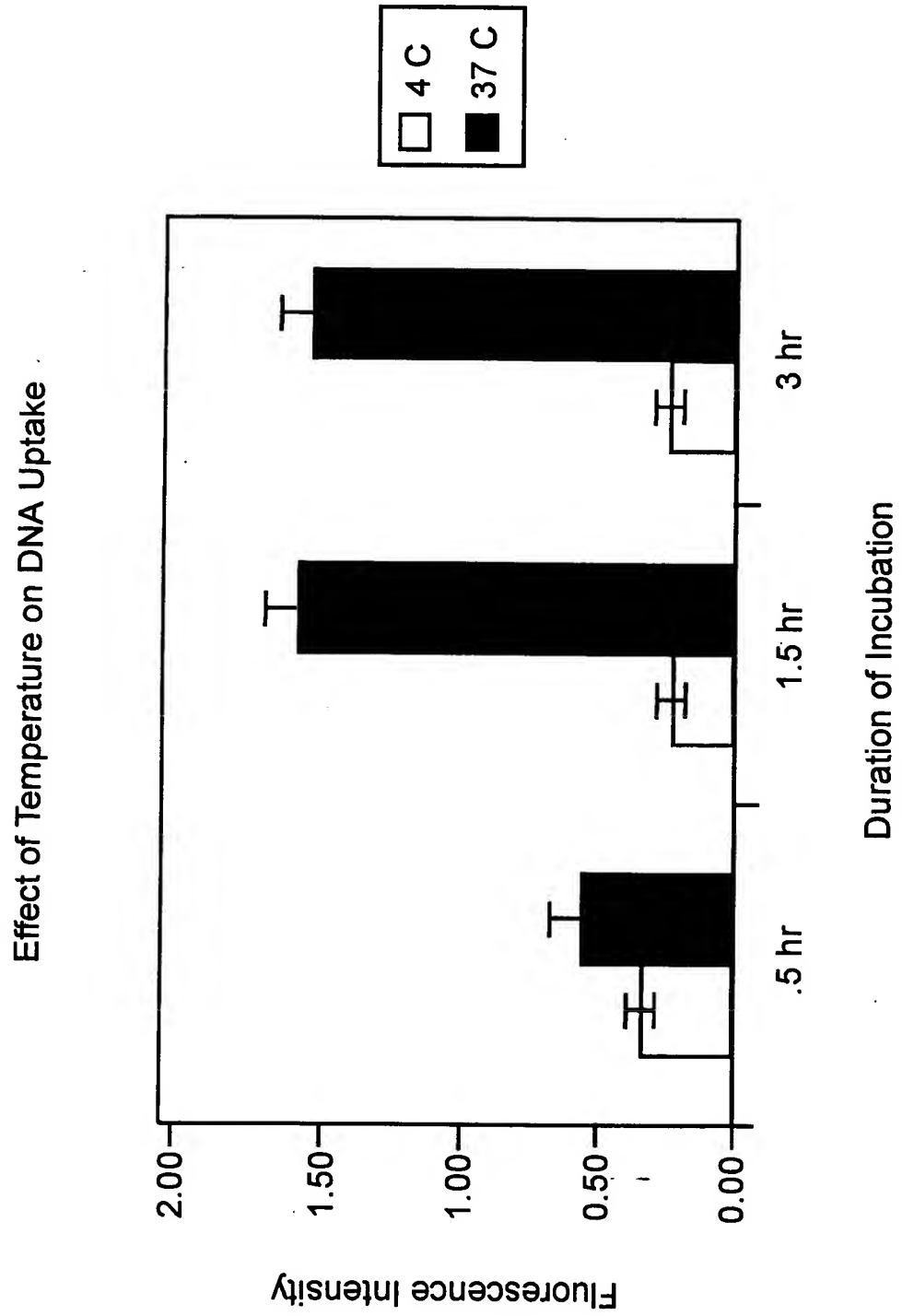
*Fig. 17A**Fig. 17B*

*Fig. 18*

*Fig. 19*

*Fig. 20*

*Fig. 21*

*Fig. 22*

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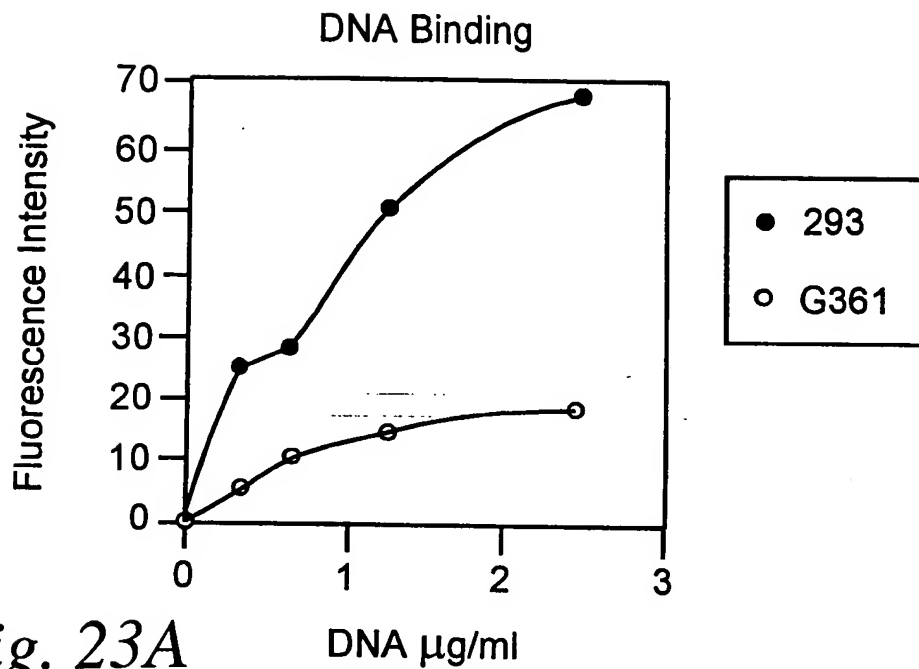


Fig. 23A

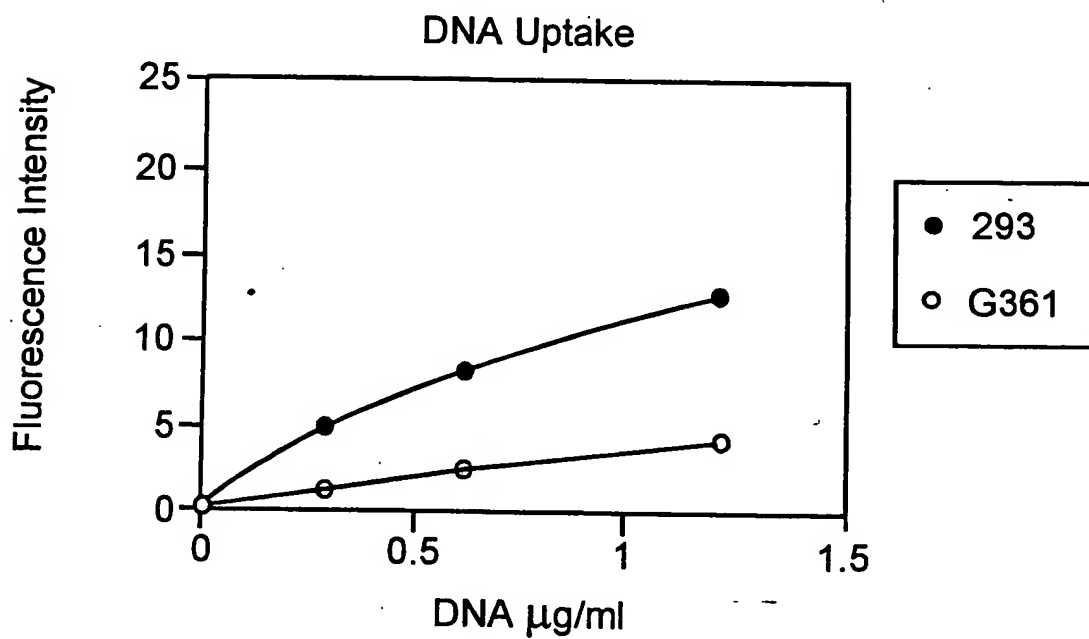
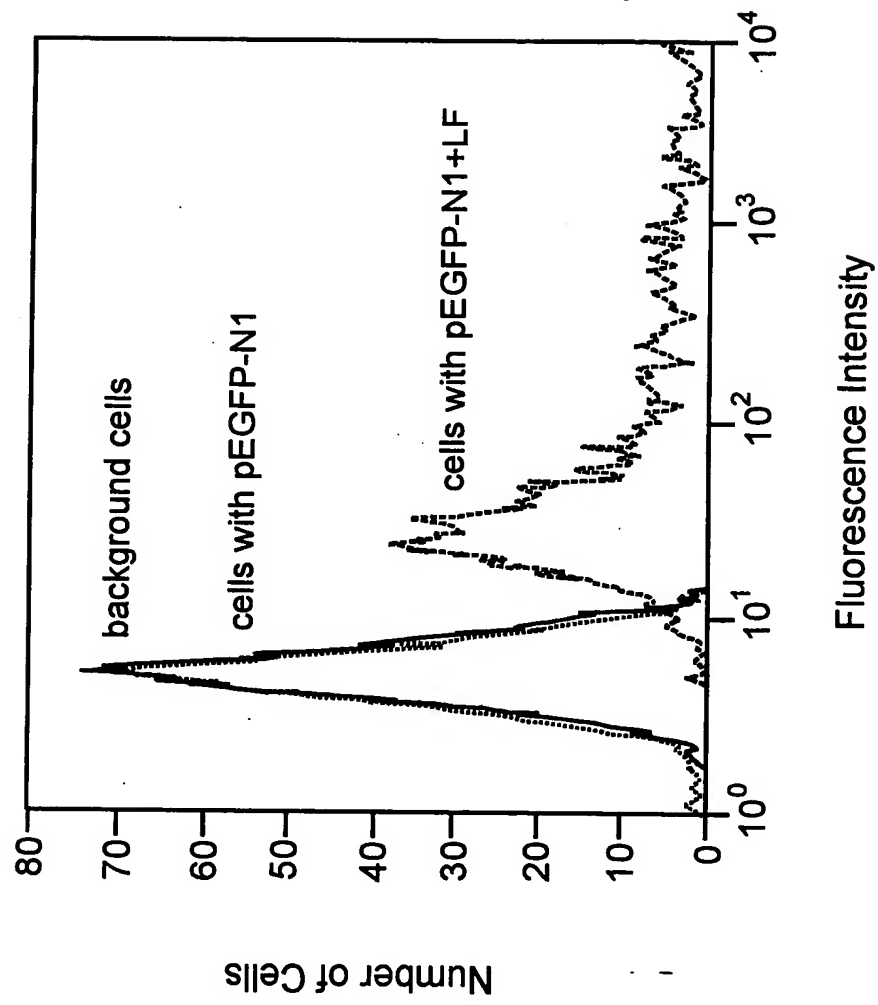
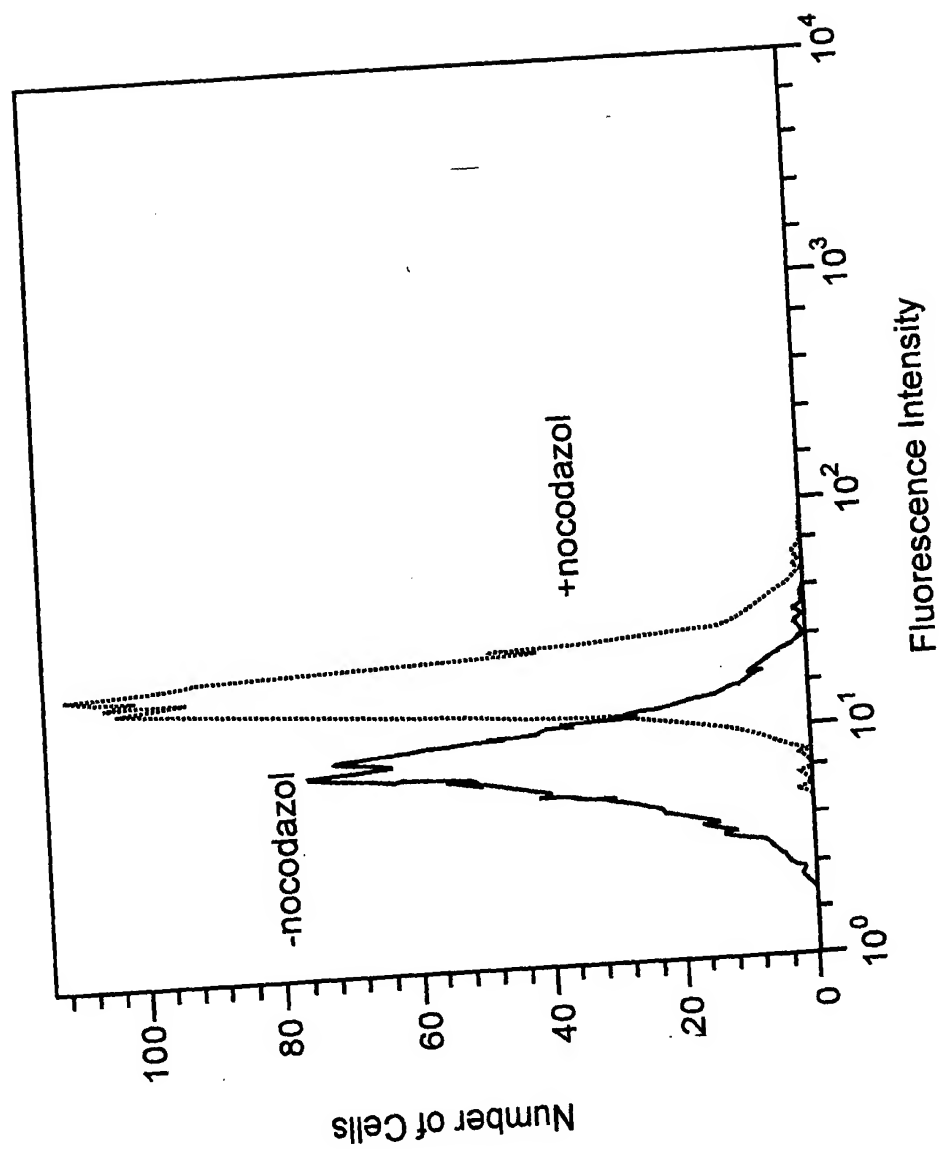


Fig. 23B

*Fig. 24*

*Fig. 25*